

RAW SEQUENCE LISTING ERROR REPORT

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK.SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
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Revised 01/29/2002



OIPE

RAW SEQUENCE LISTING DATE: 02/11/2002 PATENT APPLICATION: US/10/044,569 TIME: 09:42:29

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70

85

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40 cgg gag ttc cag ggc agc gtc acc atg acc gcg gac aca tct aca gac

41 Arg Glu Phe Gln Gly Ser Val Thr Met Thr Ala Asp Thr Ser Thr Asp

75

90

Output Set: N:\CRF3\02112002\J044569.raw

Does Not Comply
Corrected Diskette Needed

240

288

1 <110> APPLICANT: D. Collen Research Foundation 2 <120> TITLE OF INVENTION: Method and pharmaceutical composition for preventing and/or treating systemic inflammatory 4 <130> FILE REFERENCE: C1968<150> US 60/261,405<151> 2001-01-11<160> 8 W--> 5 <140 > CURRENT APPLICATION NUMBER: US/10/044,569 0 <160> NUMBER OF SEQ ID NOS: 5 <170> SOFTWARE: PatentIn version 3.1<210> 1<211> 450<212> DNA<213> Homo (sample of submitted file) Please contact Robert Wax 1 at 703-306-4119 or 703-308-4216 for assistance ERRORED SEQUENCES W--> 6 <220> FEATURE: <221> V_region<222> (1)..(450)<223> W--> 6 <220> FEATURE: <221> V_region<222> (1)..(450)<223> W--> 8 <220> FEATURE: <221> CDS<222> (1)..(450)<223> W--> 10 <220> FEATURE: <221> misc_feature<222> (130)..(159)<223> complementary determining W--> 11 region number one W--> 13 <220> FEATURE: <221> misc_feature<222> (202)..(258)<223> complementary determining W--> 14 region number two W--> 16 <220> FEATURE: <221> misc_feature<222> (343)..(375)<223> complementary determining W--> 17 region number three W--> 19 <210> SEQ ID NO: W--> 19 <211> LENGTH: W--> 19 <212> TYPE: W--> 19 <213> ORGANISM: E--> 19 <400> SEQUENCE: 1 20 atg gac tgg acc tgg agg atc ctc ttc ttg gtg gca gca gct aca ggc 48 21 Met Asp Trp Thr Trp Arg Ile Leu Phe Leu Val Ala Ala Ala Thr Gly 10 96 24 acc cac gcc cag gtc caa ctg gta cag tct ggg gct gag gtg aag aag 25 Thr His Ala Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys 28 cct qqq qcc tca qtq aaq qtc tcc tqc aaq qtt tcc qqa tac acc ctc 144 29 Pro Gly Ala Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu 32 act gaa tta ccc gtg cac tgg gtc gga cag gct cct gga aaa ggg ctt 192 33 Thr Glu Leu Pro Val His Trp Val Gly Gln Ala Pro Gly Lys Gly Leu

38 65

42